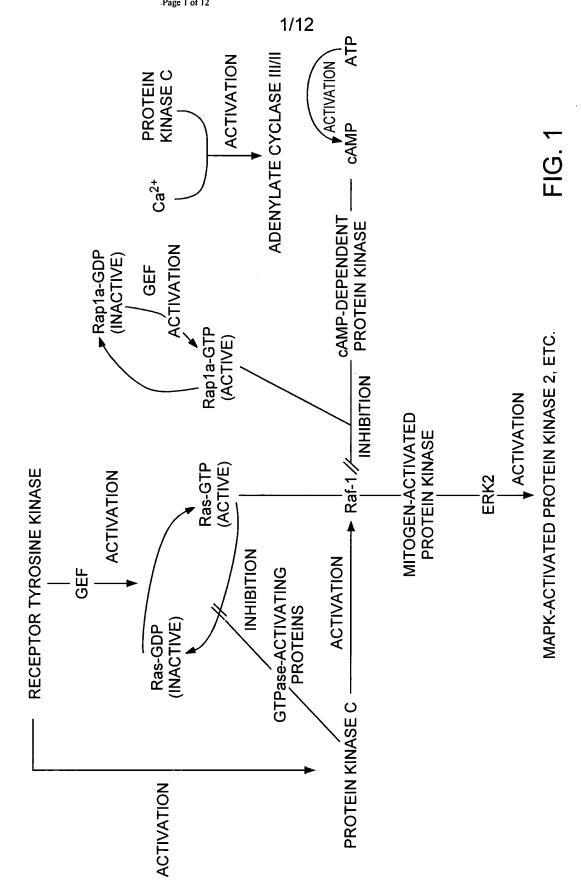
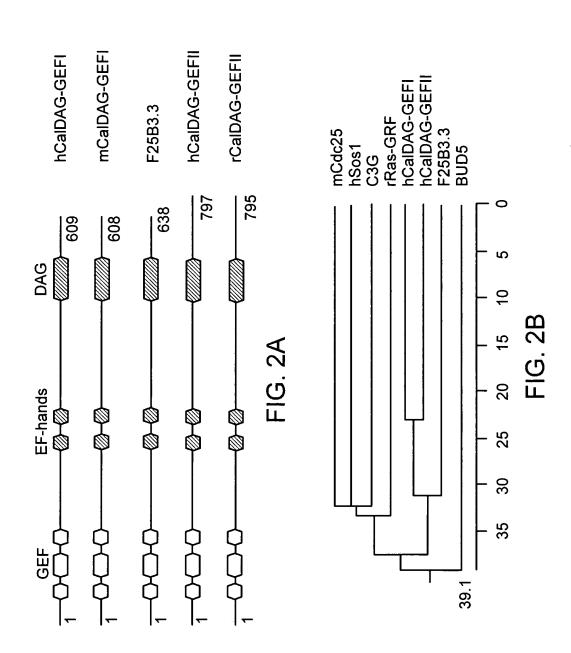
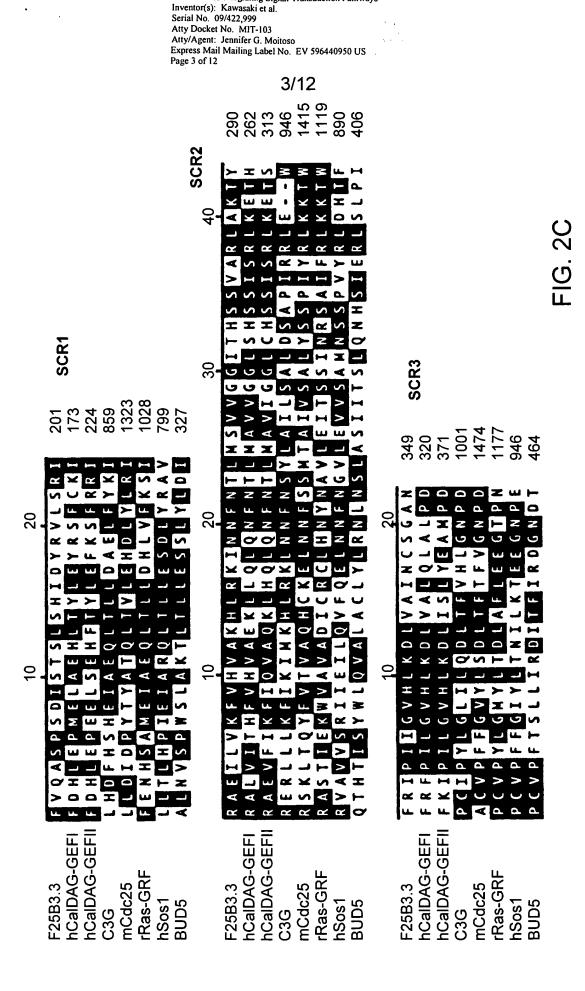


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NAGTLDLDKG NASTLDLDKG
E G R R R R R R R R R R R R R R R R R R
NSVSQUVQLNILSKP NSVSQUVQLNILSKP
H L K D L Y A L Q H L K D L Y A L Q
TSAAK
VACRHC

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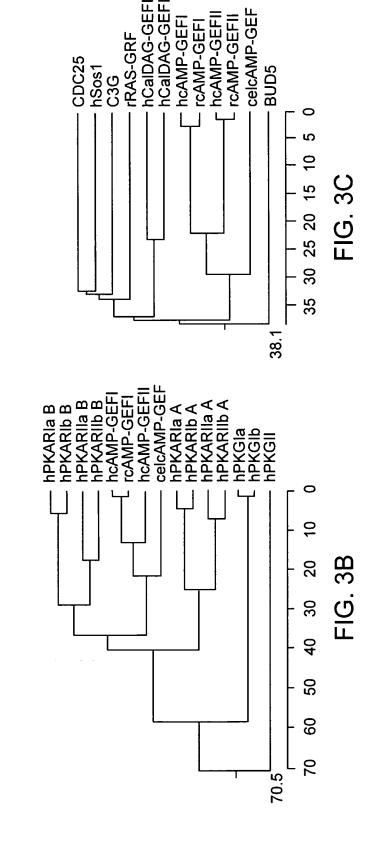
473 452 447 34 168 113 65 hCalbindin D28K hCalcineurin B hParvalbumin α hCalDAG-GEFI **Calmodulin** hTroponin C

FIG. 21

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h cAMP-GEFI

881

GEF

씸

CAMP

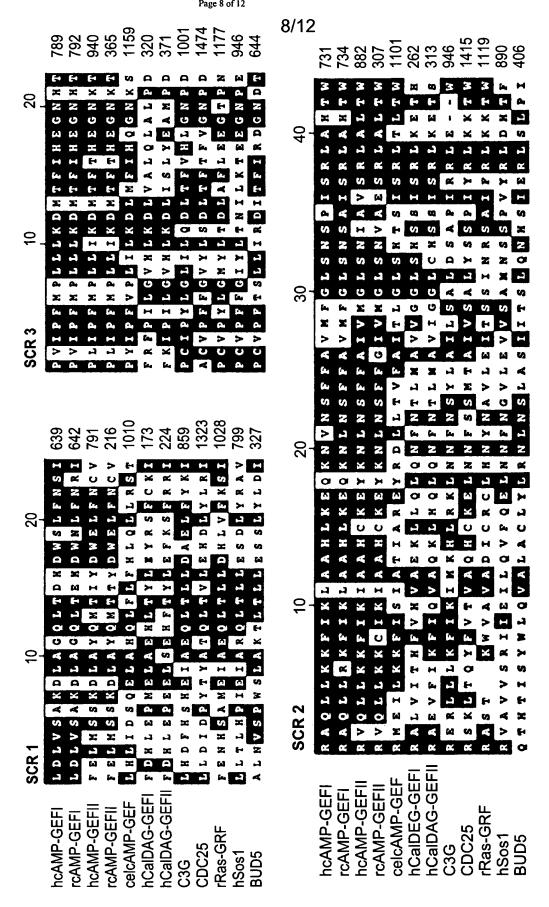


FIG. 3D

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hcAMP-GEFI hcAMP-GEFII	MVL
hcAMP-GEFI hcAMP-GEFII	YAVLAGSLDVKVSETSSHQDA
hcAMP-GEFI hcAMP-GEFII	PYGVMETGSNNDRIPDKENTP
hcAMP-GEFI	L V D G I L A L G L G V H S R S Q V V G I
hcAMP-GEFII	L V D W M M Q Q T P C V H S R T Q A V G M
hcAMP-GEFI	S Q R G P D A L L T V A L R K P P G Q R T
hcAMP-GEFII	S Q M G P D A H M R M I L R K P P G Q R T
hcAMP-GEFI	G S V N V V T H G K G L V T T L H E G D D
hcAMP-GEFII	G S V N V V I Y G K G V V C T L H E G D D
hcAMP-GEFI	A S Q G A - G P S R P P T P G R N R Y T V
hcAMP-GEFII	V P A G N R A S N Q G N S Q P Q Q K Y T V
hcAMP-GEFI	G G S E Q E R S T Y V C N K R Q Q I L R L
hcAMP-GEFII	Q G T E Q E K M D Y A L N N K R R V I R L
hcAMP-GEFI hcAMP-GEFII	S P Q M K A R N L P V W L P N Q D E P L P A P Q K K H K V L L Q Q F N T G D E R - A
hcAMP-GEFI hcAMP-GEFII	G D A I G L Q P D A R G V A T S L G L N E G E K V V L K P N D V S V F T T L T I N G
hcAMP-GEFI	HYVLGPQHLRDVTTANLERFM
hcAMP-GEFII	YHTFGRHNFKK-TTANLDLFL
hcAMP-GEFI hcAMP-GEFII	R L A H T W E R L P H K V R K L Y S A L E R L A L T W E K L P S K F K K F Y A E F E
hcAMP-GEFI	AARMLHHCRSHNPVPLSPLRS
hcAMP-GEFII	TARTVRYYRSQPFNP

FIG. 3F-1

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RRMHRPRS
S C I Q G
LTNSEESLDFSESLEQASTERVLRAGRQLHQHL LIEPHVPLRPANTITKVPSEKILRAGKILRNAI
CQVLLDEGALCHVKHDWAFQDRDAQFYRFPGPE WQVLLEDGVLNHVDQEHHFQDK-YLFYRFLDDE
DEELDLIFEELLHIKAVAHLSNSVKRELAAVLL VDDLEIIYEELLHIKALSHLSTTVKRELAGVLI
F G Q L A L V N D A P R A A T I I L R E Y N C H F L R V D K Q D F F G K L A L V N D A P R A A S I V L R E D N C H F L R V D K E D F
M S G T P D K I L E L L L E A M G L D S S A H D P K E T F L S D F M S G T P E K I L E H F L E T I R L E A T L N E A T D S V L N D F
V S Q W V A L Y G S M L H T D P V A T S F L H K L S D L V G R D T V L Q W A A M Y G D L L Q E D D V S M A F L E E F Y V S V S D D A
GSSCAIQVGDKVPYDICRPDHSVLTLQLPVTAS QKRQPIRGSDEVLFKVYCMDHTYTTIRVPVATS
R L F V V N P Q E V H E L I P H P D Q L G P T V G S A E G L D L V R L F A C P R E Q F D S L T P L P E Q E G P T V G T V G T F E L M
R R F N E L Q Y W V A T E L C L C P V P G P R A Q L L K K F I K L R R F N E I Q F W V V T E I C L C S Q L S K R V Q L L K K F I K I
R L L D P S W N H R V Y R L A L A K L S P P V I P F M P L L L K D S L M D P S R N H R A Y R L T V A K L E P P L I P F M P L L I K D
RVSHLHEDSQVARISTCSEQSLSTRSPASTWAY

FIG. 3F-2

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CSYQLLLE	20 80
L L R I E Q K D F K A L W E K Y R Q Y M A G L L A P	32 160
LATCPNLIRDRKYHLRLYRQCCSGRE	92
LSRAPHMIRDRKYHLKTYRQCCVGTE	240
PE PVGTHEM EEELAEAVALL HEDAPLPTEEEKKECDEELQDTMLLL	166 319
FEPHSKAGTVLFSQGDKGTSWYIIWK	246
FESHAKGGTVLFNQGEEGTSWYIILK	399
N R I I K D V E A K T M R L E E H G K V V L V L E R	326
N R I L R D V E A N T V R L K E H D Q D V L V L E K	479
LLTHRVFMPSAQLCAALLHHFHVEPA	405
IMMHCVFMPNTQLCPALVAHYHAQPS	559
R L S N L L R E Q W P E R R R C H R L E N G C G N A R M I A A L K E Q L P E L E K I V K Q I S E - D A K	485 638
VREVMAALAQEDGWTKGQVLVKVNSA	565
VKEVISAVADKLGSGEGLIIVKMSSG	717
SAKDLAGQLTDHDWSLFNSIHQVELI	645
SSKDLAYQMTIYDWELFNCVHELELI	797
A A H L K E Q K N V N S F F A V M F G L S N S P I S	725
A A H C K E Y K N L N S F F A I V M G L S N I A V S	876
MTFIHEGNHTLVENLINFEKMRMMAR	805
MTFTHEGNKTFIDNLVNFEKMRMIAN	956
VQQLKVIDNQRELSRLSRELEP	881
VRQLNVIDNQRTLSQMSHRLEPRRP	1011